

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/553,884

Source: Pg 10

Date Processed by STIC: 11/2/05

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PCT

RAW SEQUENCE LISTING

DATE: 11/02/2005

PATENT APPLICATION: US/10/553,884

TIME: 09:44:23

Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF4\11022005\J553884.raw

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3 <110> APPLICANT: Agrinomics LLC
5 <120> TITLE OF INVENTION: GENERATION OF PLANTS WITH ALTERED OIL CONTENT
7 <130> FILE REFERENCE: 6616-72014-06
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/553,884
C--> 9 <141> CURRENT FILING DATE: 2005-10-20
9 <150> PRIOR APPLICATION NUMBER: PCT/US2004/012278
10 <151> PRIOR FILING DATE: 2004-04-19
12 <150> PRIOR APPLICATION NUMBER: 60/464,558
13 <151> PRIOR FILING DATE: 2003-04-22
15 <160> NUMBER OF SEQ ID NOS: 6
17 <170> SOFTWARE: PatentIn version 3.2
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 693
21 <212> TYPE: DNA
22 <213> ORGANISM: Arabidopsis thaliana
24 <400> SEQUENCE: 1
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27 cacaggtatt tcgttgaaca gcttatgtgt gcttcttcct tggatgaaac taatcgtatc      120
29 tctctcgaag aacaacaatc tctcgtggcc cagtttctat ctcactgtct tcaatactac      180
31 caagagaaat tcgcctccgt ttccctcgcc ggggacaacg ttttcacttt cttctgccca      240
33 ccgtggttta actcctacgc taaacttatt ttatgggtcg gcgatttcaa gccttctctt      300
35 gtgtttaaac tcaccgaggt ctccgtggcc gacctcacgc gccaccagaa agaccggatc      360
37 tcgagtctta agtcggagac taggaggaaa gagagagaag ttatgcgaga ttctgccttc      420
39 gtgcaacaaa gcgtggcgga tccgccggtg atgctcgcgg cgaggcgcggt gggagcggtg      480
41 ggaatggtgg acggagaaga aacggatttg gaggaggcga tggagggtgct taaagctggg      540
43 atggcggcag cgatgaacaa cgctgatcag ctacggtgtt cgacggtggg gaaagtgggtg      600
45 gagattctta ctccgccgca agcgattaaa gtgttgagga caatcggaca gcttcacctc      660
47 cgtctgagag acagagacca agaaagagct taa                                693
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51 <211> LENGTH: 230
52 <212> TYPE: PRT
53 <213> ORGANISM: Arabidopsis thaliana
55 <400> SEQUENCE: 2
57 Met Pro Asn Thr Ser Ser Ser Gln Ser Phe Thr Ile Phe Val Asp Gly
58 1          5          10          15
61 Trp Leu Ile Arg His Arg Tyr Phe Val Glu Gln Leu Met Cys Ala Ser
62          20          25          30
65 Ser Leu Asp Glu Thr Asn Arg Ile Ser Leu Glu Glu Gln Gln Ser Leu
66          35          40          45
69 Val Ala Gln Phe Leu Ser His Cys Leu Gln Tyr Tyr Gln Glu Lys Phe
70          50          55          60
73 Ala Ser Val Ser Leu Ala Gly Asp Asn Val Phe Thr Phe Phe Cys Pro
74 65          70          75          80

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77 Pro Trp Phe Asn Ser Tyr Ala Lys Leu Ile Leu Trp Val Gly Asp Phe
78                               85                               90                               95
81 Lys Pro Ser Leu Val Phe Lys Leu Thr Glu Val Ser Val Ala Asp Leu
82                               100                              105                              110
85 Thr Arg His Gln Lys Asp Arg Ile Ser Ser Leu Lys Ser Glu Thr Arg
86                               115                              120                              125
89 Arg Lys Glu Arg Glu Val Met Arg Asp Phe Ala Leu Val Gln Gln Ser
90                               130                              135                              140
93 Val Ala Asp Pro Pro Val Met Leu Ala Ala Arg Arg Val Gly Ala Val
94 145                               150                              155                              160
97 Gly Met Val Asp Gly Glu Glu Thr Asp Leu Glu Glu Ala Met Glu Val
98                               165                              170                              175
101 Leu Lys Ala Gly Met Ala Ala Ala Met Asn Asn Ala Asp Gln Leu Arg
102                               180                              185                              190
105 Cys Ser Thr Val Gly Lys Val Val Glu Ile Leu Thr Pro Pro Gln Ala
106                               195                              200                              205
109 Ile Lys Val Leu Arg Thr Ile Gly Gln Leu His Leu Arg Leu Arg Asp
110                               210                              215                              220
113 Arg Asp Gln Glu Arg Ala
114 225                               230
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118 <211> LENGTH: 1163
119 <212> TYPE: DNA
120 <213> ORGANISM: Arabidopsis thaliana
122 <400> SEQUENCE: 3
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125 gacactgatg agaggaacca gatgtttgaa caaggacttg ttgctgtccc cacagcttct      120
127 gattctagtg acaaatcaag ggacaaacta gatcagaaga cacttcggcg tcttgcccaa      180
129 aatcgtgaag ctgcccggaa aagccgttta cgaaagaagg catacatcca aaaccttgag      240
131 agtagcagat tgaaacttac tcagttagag caagagcttc accagactcg tcaacagggt      300
133 atttttatatt ctacatcagg agatcaacct caatcaacaa gcggaaatgg agctttggca      360
135 tttgacatgg agtatgcacg ctggttggaa gagcacaaca aacatgtaaa tgagttgagg      420
137 cttgcagtca atgcacatgc cggtgataat gatctccgtg gtattgttgg tagtgttatg      480
139 gcacactacg atgaattttt caggctcaag ggtgtggcag ctagatcaga tgtttttcat      540
141 gtgctgtctg ggatgtggaa gaccctgct gagagatgtt tcatgtggtt aggtggcttc      600
143 cgatcatctg aggttcttaa gttactggca ggtcacctag agcctcttac tgatcagcag      660
145 cttgttggtg tatccaacct gcagcagtc tcccaacaag ctgaagatgc tctttctcaa      720
147 gggatggaag cattacaaca gtcgcttgca gaaactctag catctggatc cctgggccct      780
149 gctggacctt ctggcaatgt tgcaaattac atgggacaaa tggcgatggc tatgggaaaa      840
151 cttggcaccc tagagaactt tctacgacag gctgataatc tacggctgca aacacttcaa      900
153 caaatgcaac gcatcttaac caccgcacaa tccgcacgag ccctacttgc aataagcgac      960
155 tacttctctc ggctgcgtgc tttgagttct ctttggcttg cccgtccaag ggaataaatt     1020
157 caattaactc ataaatagga ttttttttgt actgtctcta gtatgtcaat tagtttcttc     1080
159 acattgctca tgtgttgatg ggatatttgg atctaagatc ttatgaatgg atgacatgta     1140
161 tgtgttcaag tgaaaaaaaa aaa                                     1163
164 <210> SEQ ID NO: 4
165 <211> LENGTH: 1024
166 <212> TYPE: DNA
167 <213> ORGANISM: Arabidopsis thaliana

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170 <220> FEATURE:

171 <221> NAME/KEY: misc_feature

172 <222> LOCATION: (416)..(417)

173 <223> OTHER INFORMATION: n is a, c, g, or t

175 <400> SEQUENCE: 4

W--> 176 tggaaagaac ctttttttgg gttgggggggt tcaaacctgg aatggccttt caggttgtga 60
 178 acgcagcttt ggagggtttg tcagaacagc aaaaggagag gctgagtttg ctcaaccaag 120
 180 agaccaaggt gaaggagaga gccctcaatg atgagttggc caagcttcat gagagtgtgg 180
 182 cagctccacc gctggtggac atggccagga gccatggtag ggtgtgtttc agtaggtcct 240
 184 tcatggcaga tgggggttct tccgttccaa gcactttcag atagacattg gagaatctgg 300
 186 aggcaaatgc agatgctttg aggacaaaca catctttgaa gattctccag atactgagac 360
 188 cttctcagct tgtttcattc ttggctgctg tggctgagct tcagatcagg attggnnctt 420
 190 ggggttttga caaggatgcc ctgaatggag gccaaagggtg aaagtcaatt aacgggaatt 480
 192 tgaaggcttt tttatgtaag ttgctttcag tttttgtctt ttttgggaac tggtttgagt 540
 194 tggaggatgg atacataggc atttagcctt ttagcatcat gatgatggat ggtgtgcatt 600
 196 gggcagtgg catgttcact gcctgtaatg ttgtttctct ctacctggtt tgggtgggcat 660
 198 tgaacaatga tgaacaaaac aaggccatgc aatgcatact tttatctaga ggctaaaagc 720
 200 atcaatatat ttgtacctct tgagatagag gcaagtactc caggttctct ctactcaagc 780
 202 agctacccca gatttcaatt gggtgctata tttaccatag taagtaggta gtagatacct 840
 204 aaggattatt tatttttctt ttttttgtga tattaagaca tgttttctaa tttctagtga 900
 206 agaacttggg tggatgtata ggatgtggta ccttgaaaat tacttttcaa cgtcccactc 960
 208 actttgacat tgcacaaag taagtttctc cctaaggggt ctttttggtc ctttcctcaa 1020
 210 caca 1024

213 <210> SEQ ID NO: 5

214 <211> LENGTH: 1234

215 <212> TYPE: DNA

216 <213> ORGANISM: Arabidopsis thaliana

218 <400> SEQUENCE: 5

219 ggcgcagtgt gcaggtgcaa ctgagcaatt gcagacctgg gagttgcttc cattatcttg 60
 221 cacacactac taggtggcta gccatcgagc acaatcatgt ccaagctggc aagtagtgct 120
 223 tcataccttg tctccacctc gccaaacggt tctccttccc gtgaaacgtt tcgcaaattc 180
 225 ttcgagtgtt ggcttggaga gcaaaacaat tatctcgaac aactcatctc aacctgtaaa 240
 227 gattatgatc acaacagaaa aaactcccc cagtcactctc aggcaaccct ccagcctctt 300
 229 atcaaccgtg ttcttgagca ttatgaacat tattatagat ccaagtcaag atgggcaaaa 360
 231 gatgatgtgt tatccatgct ctctccttca tggaccagca ctctcgagca tgcttttctt 420
 233 tggatcggtg ggtggcggcc ctctgtggct tttcacttgc tctactcaa gtcaggtcat 480
 235 cagtttgagg ctcaactcca tgagttgatt tgtgggttgg ggacaggtga cttgggtgac 540
 237 ctttcagcta gtcaactcac ccgagttgat cagttacaga ggaagactgt caggggaagag 600
 239 aacgagctga ctgagaagct tgcaaagcaa caagaaactg tagcagactc gtctatggtg 660
 241 gagttagcac atgaggtgac tgagttgttg aggagcgaga acacgggtga tgaagtggag 720
 243 gaagagcgag ttgagtcaac tctggcacct aaaaaggatg gattgcagga aatcttgag 780
 245 atggctgatg atctacggct gagaactatc aaaggtgtta tcgagathtt gactccaatc 840
 247 caagccgttc atttcttgat tgcagctgct gagttgcact tgcgccttca tgactggggc 900
 249 aagaagggtg attgggcacg ccgcgtccac cactgatatg cgatttatta ggggataatc 960
 251 atccatgaat tacaccata ttcgctgttc tgttgacata attaagaaat ggggagactg 1020
 253 attttggttt tagaggcagg ataagcaaga tacgcagtaa tgtttaatta tattttcaga 1080
 255 ttgaatctat ctatgtttac tatgtattat ccatcaggtg ctttgagtcc tcaagcactg 1140
 257 gcattttagt ttaatatatg tattatgtat gatgttgcca gttaatttct ccaacttggc 1200
 259 tttggctata attgaagtat tttgaaagat tgaa 1234

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262 <210> SEQ ID NO: 6
263 <211> LENGTH: 1083
264 <212> TYPE: DNA
265 <213> ORGANISM: Arabidopsis thaliana
267 <400> SEQUENCE: 6
268 gatagtctct atttatattg tagatatata aaccaactct ccatgatgag tactagcaaa      60
270 aatgggctag aaaatggcaa atcattccac aagttttttt gaatcatggc tcgttaaaca      120
272 aaatcaagat ttggatcagc ttgtacgtgc ctcaaaagac gacgacaaca acaacaagaa      180
274 caacaatgac atgatgttgt catctttaat tcatagtgtg gtgaaacatt atgaagaata      240
276 ttatagagag aaatcacgat acgctattag tgatatttta ggcatgttgc acccctcatg      300
278 gttaagtaat cttgaagatg catttttatg gattgggtgga tggagaccta gtatggcttt      360
280 tcatttgtaa tactcaaaat caggtatata acttgaagct aatcttcatg agttaattag      420
282 aggatttaac acaaaagatt taggaaattt aagtggtaat caacttgtat taattgatga      480
284 gttacaacat aagacaataa gtgaagaaag aaagctaagt gaaaatttag ctaaagttca      540
286 agaaacttta gctgatgcat caatggttga attatcacat gttgtgagtg aattaatgag      600
288 ggatgatcaa ttagttgtta atgatgaaga ggaaaaaatt aagaaaaata ttagtaaaaa      660
290 agaggaaagt ttattggatt tgttgaaaaa agctgatgat ttaagggtta gtacaattaa      720
292 agagattttg agaattttga catcaacaca aggtgttcat ttcttgattg ctgctgctga      780
294 acttcatttg aggattcatg aatggggtaa gaagaaagat gctgctgttt ctcatacata      840
296 atgggtcatgt caaacaataa acgacgagcc aaattctgag gggacgtctg tcagaaatta      900
298 gtcattcaga tctattagaa tgatacagag aagatcaacg tggacttaa tagttatgaa      960
300 tattcttaat tactacttaa ggagctagcc aaatgttatt agtattaatt actaggtaga     1020
302 ttgtaatagg ataaagttct tttaagggct tttttttttg tttcattttt gtttatgtag     1080
304 tga                                           1083

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/553,884

DATE: 11/02/2005
TIME: 09:44:24

Input Set : A:\Sequence Listing.txt
Output Set: N:\CRF4\11022005\J553884.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:4; N Pos. 416-417

VERIFICATION SUMMARY

DATE: 11/02/2005

PATENT APPLICATION: US/10/553,884

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Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF4\11022005\J553884.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:188 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:360